

TGTCAAACACACACATAACACATAAGTGACCGTGAGTCATTAAATTTATATATATTCATCAATC

TAATCAAACATATGGAGAAGAAATCACTAGCTGGCTTATGCTTCCTCTTCTTGGTTCTCTTTGTT
M E K K S L A G L C F L F L V L F V

GCACAAGAAATTATGGTGACAGAAGCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGAC
E Q E I M V T E A R T C E N L A D K Y R G P
Δ

CATGCTTTAGTGGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGTAG
C F S G C D T H C T T K E N A V S G R C R

GGACGACTTCCGCTGCTGGTGTACTAAAAGATGTTAAATGGATCTCCTCCAACATCAAGATGTG
D D F R C W C T K R C *

CAAATAGTCTTTATAATAAACTAAATAAATAAAATGCACGCAGTATAGCTACAACCTCATCTA

TTATATGTACTCAATATCGNGCATAACGTATTAGTTATGCACTTCTATCATATGGAATAAACAT

AATAAGTAATTTTCGTNTCCAAAAAAAAAAAAAAAAAAAAA

FIG. 1

[illegible]

FIG. 2

AlfaAFP2	TGTCAAACACACACATAACACATAAGTGACCGTGAGTCATTAAATTTATA
AlfaAFP1	-----
AlfaAFP2	TATATTTCATCAATCTAATCAAACCTATGGAGAAGAAATCACTAGCTGGCTTA
AlfaAFP1	-----CAGGCTTA
	* * * * *
AlfaAFP2	TGCTTCCTCTTCCTCGTTCTCTTTGTTGAACAAGAAATTATGGTGACCGAG
AlfaAFP1	TGCTTCCTCTCTCTTGGTTCTCTTTGTTGCACAAGAAATTGTGGTGACAGAA
	***** * * * * *
AlfaAFP2	GCAGCTACTTGTGAGAATTTGGCTAACACATACAGGGGACCATGCTTCGGT
AlfaAFP1	GCCAGAACATGTGAGAATTTGGCAGATAAATATAGGGGACCATGCTTTAGT
	** * * * * *
AlfaAFP2	GGTTGTGACTTTCCTGCAAAACCAAAGAACACTTACTTAGCGGXAGGTGC
AlfaAFP1	GGTTGTGACACTCACTGCACAACCAAAGAGAACGCAGTTAGTGGAAGGTGT
	***** * * * * *
AlfaAFP2	AGGGACGACTTCCGCTGCTGCTGGATCC
AlfaAFP1	AGGGACGACTTCCGCTGCTGCTGGATCC

FIG. 3

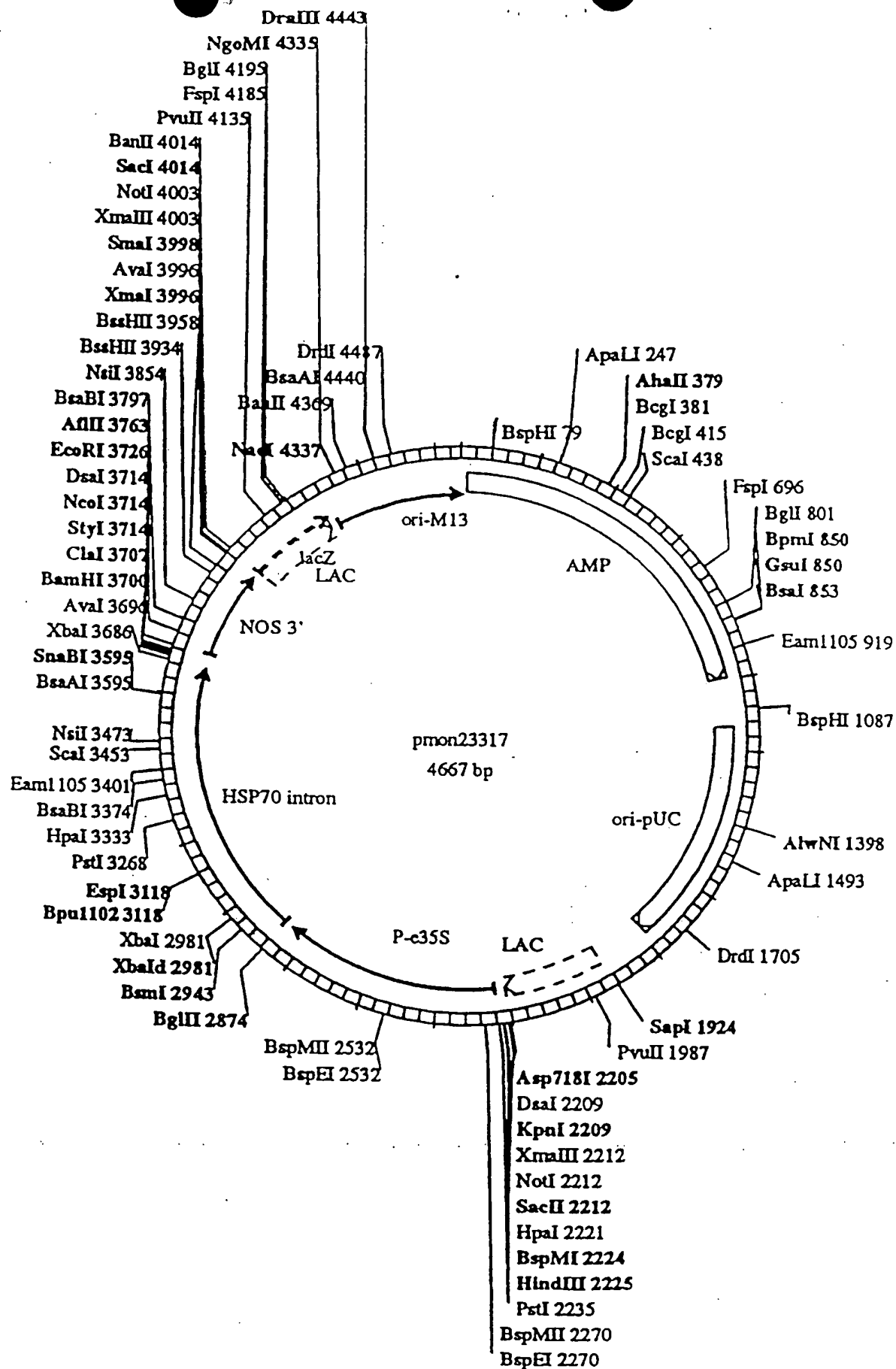


FIG. 4

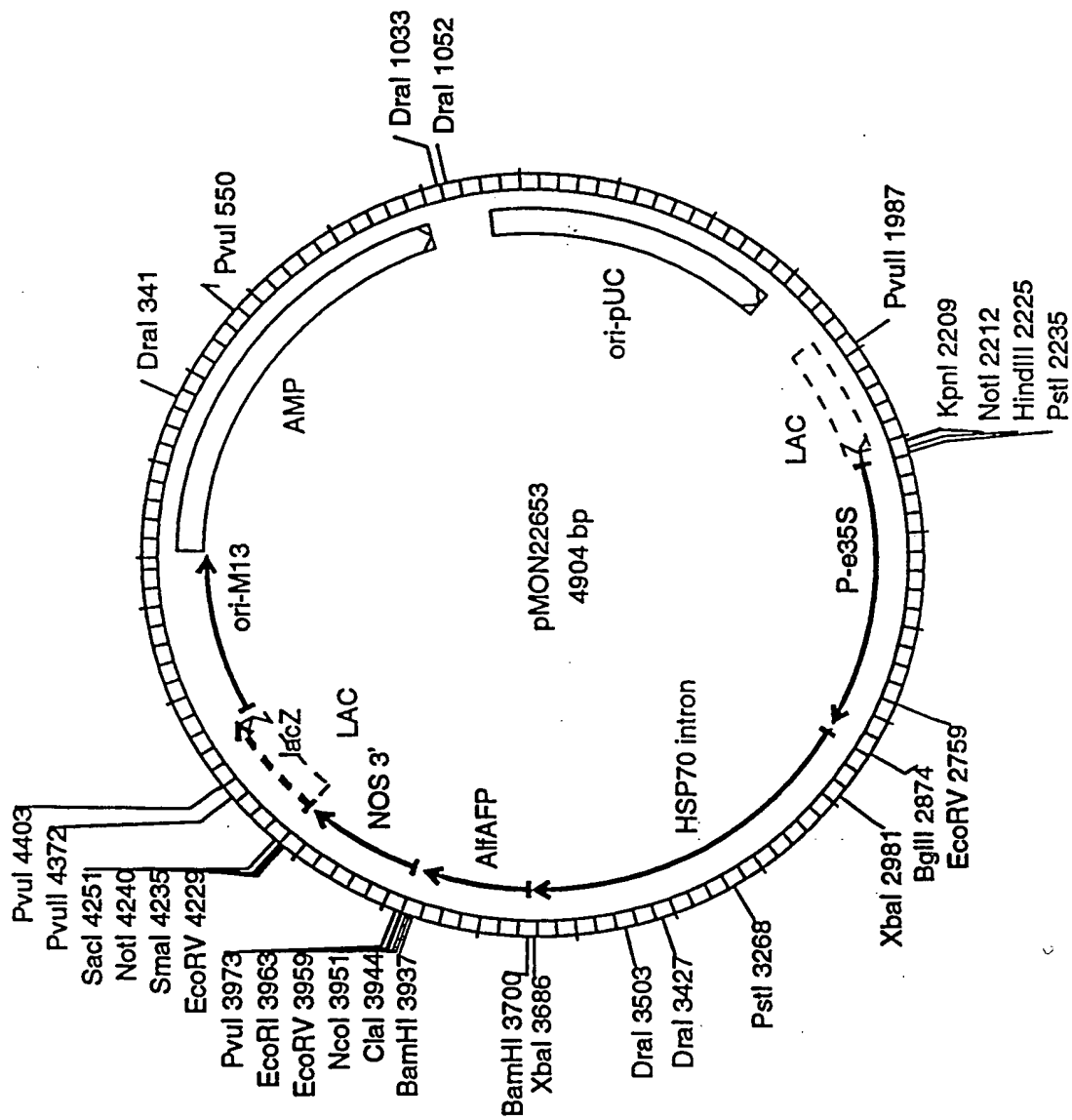


FIG. 5

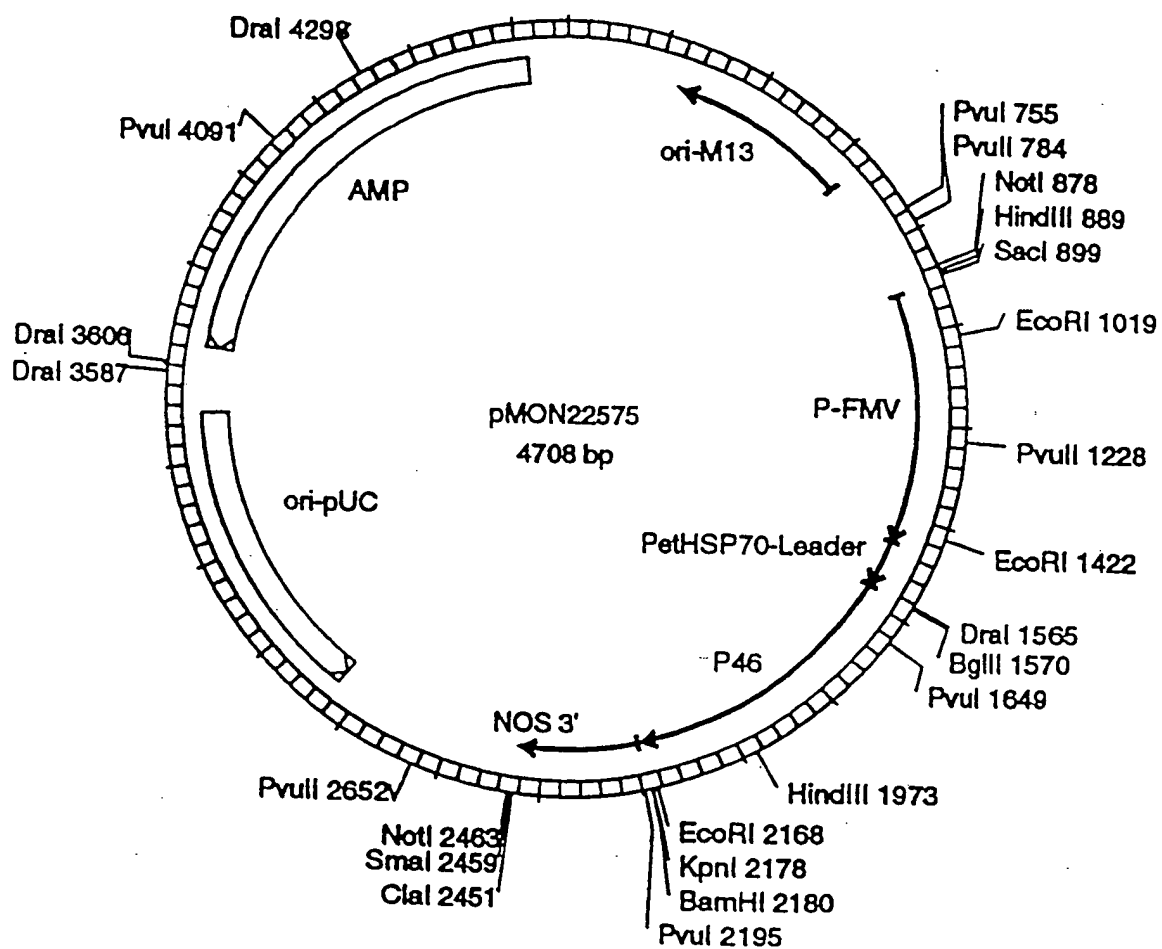


FIG. 6

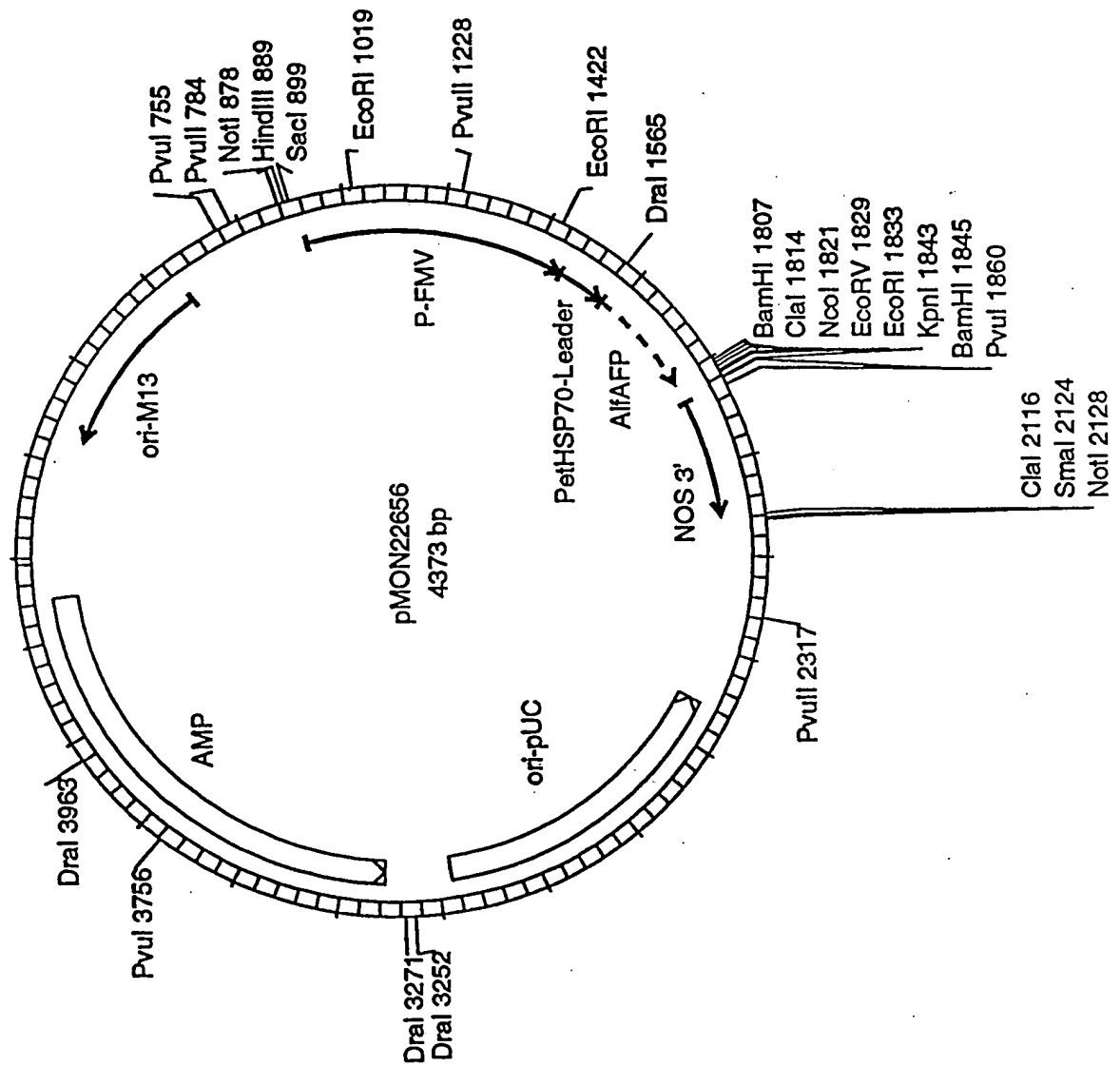


FIG. 7

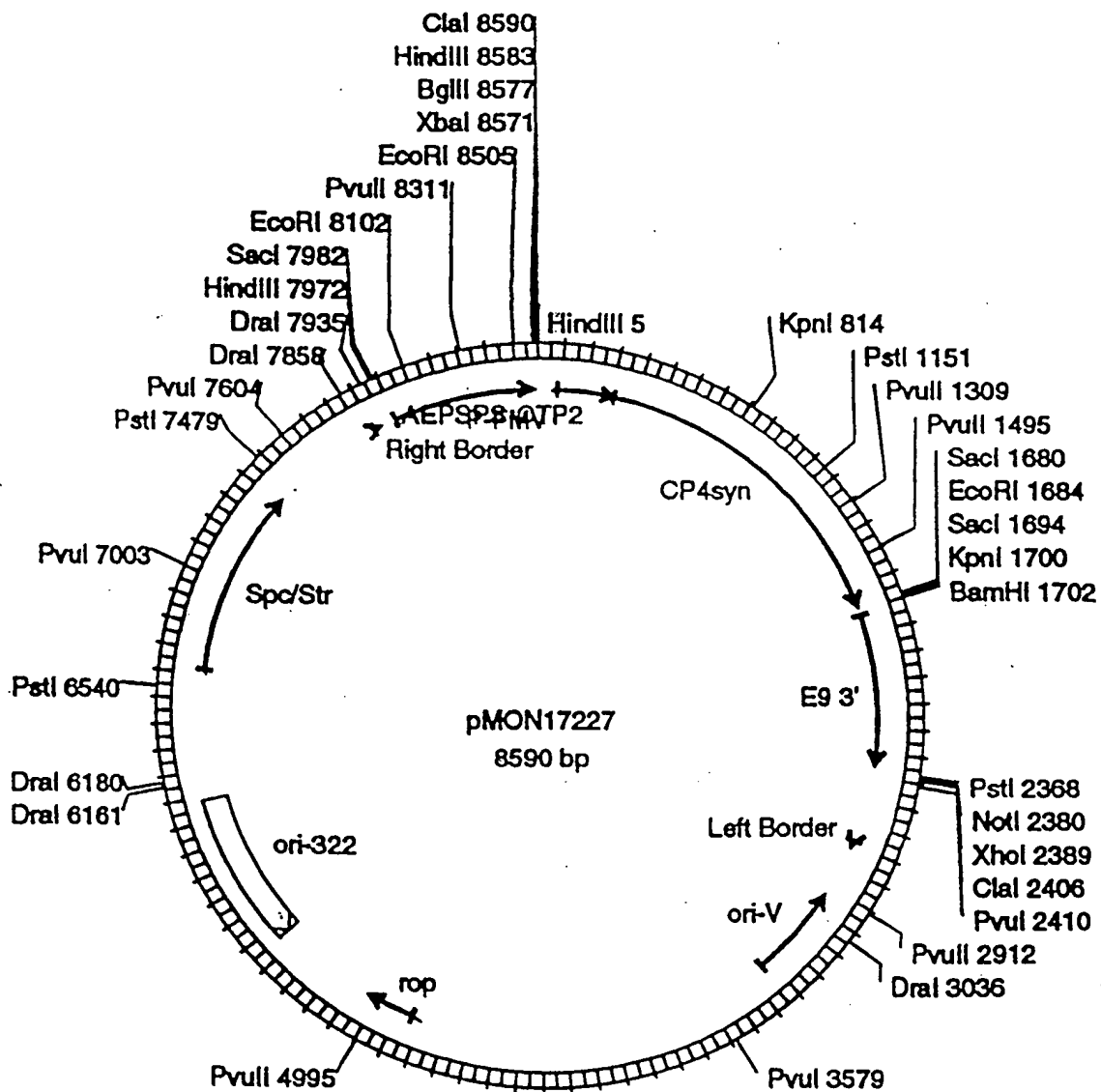


FIG. 8

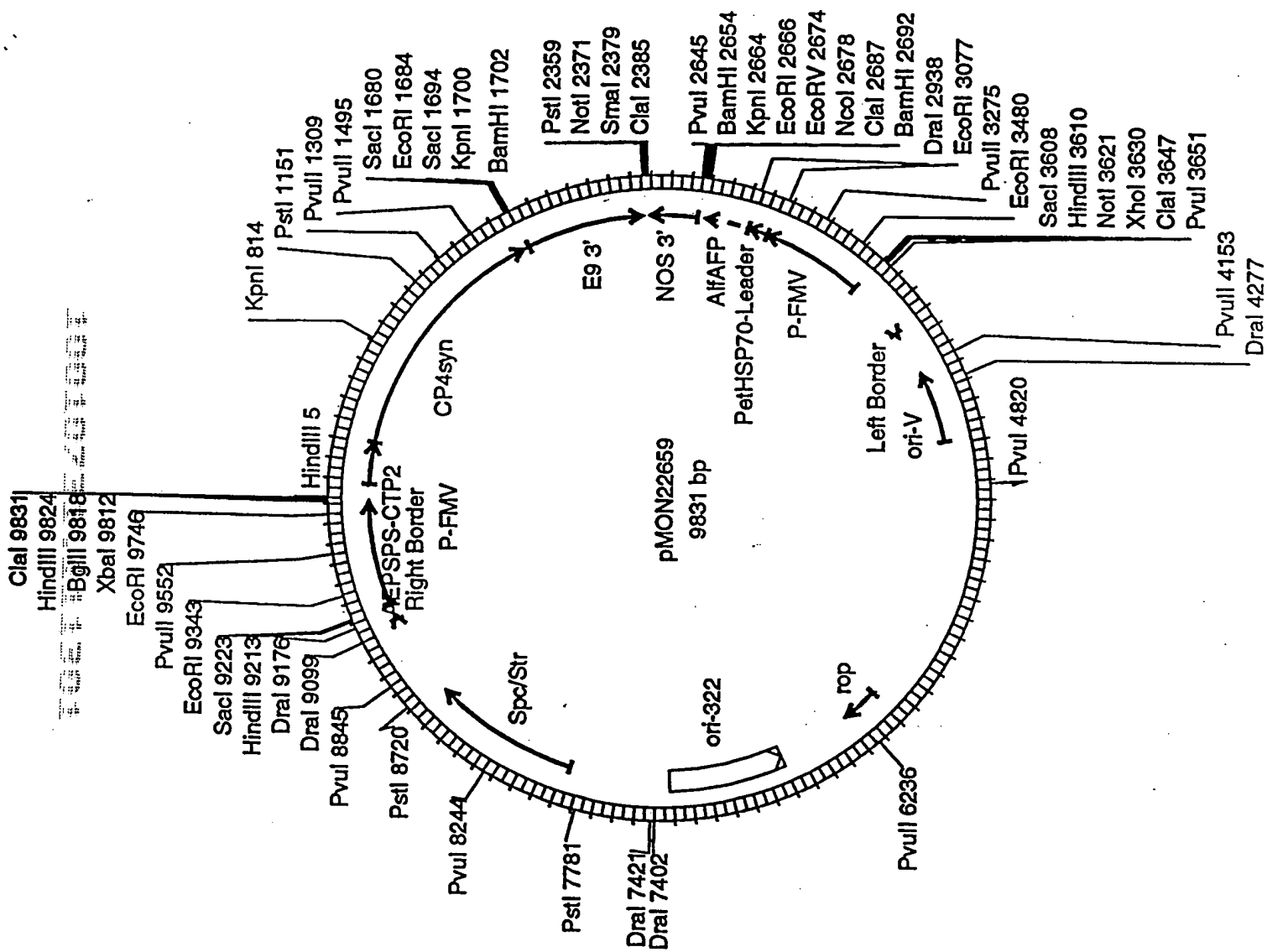


FIG. 9

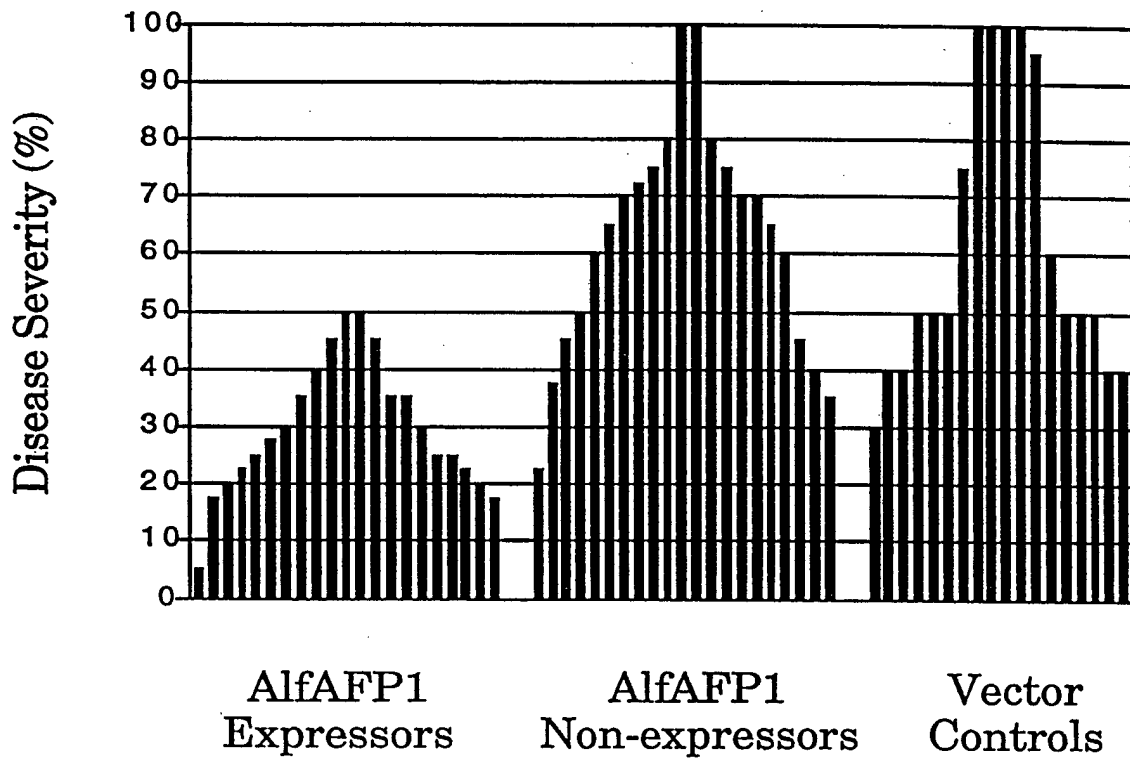


FIG. 10